

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/573,885
Source: IFwd
Date Processed by STIC: 4/17/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

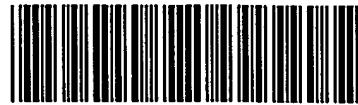
Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450**
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314**

Revised 01/10/06



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/573,885

DATE: 04/17/2006
TIME: 12:27:23

Input Set : E:\seq_SU0411.txt
Output Set: N:\CRF4\04172006\J573885.raw

3 <110> APPLICANT: SUNTORY LIMITED
5 <120> TITLE OF INVENTION: A gene encoding an enzyme for catalyzing biosynthesis of lignan,
6 and use thereof
8 <130> FILE REFERENCE: SU0411
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/573,885
C--> 11 <141> CURRENT FILING DATE: 2006-03-29
13 <150> PRIOR APPLICATION NUMBER: JP 2003-341313
14 <151> PRIOR FILING DATE: 2003-09-30
16 <150> PRIOR APPLICATION NUMBER: JP 2003-432383
17 <151> PRIOR FILING DATE: 2003-12-26
19 <160> NUMBER OF SEQ ID NOS: 79
21 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

1770 <210> SEQ ID NO: 77
1771 <211> LENGTH: 2815
1772 <212> TYPE: DNA
1773 <213> ORGANISM: Sesamum radiatum
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1797 ttctttggccg aaaaatgtct ttgttagagt cttgtgcattt gttttctata tgccaaatata 1320

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/573,885

DATE: 04/17/2006
TIME: 12:27:24

Input Set : E:\seq_SU0411.txt
Output Set: N:\CRF4\04172006\J573885.raw

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 1822 agcaatttct catcaactca taacgtactc aaataaacac gcttttcag ccaac 2815

Explanation
in
2207-2237
section
(see p.3)

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/10/573,885

DATE: 04/17/2006
TIME: 12:27:25

err
Explanation
Input Set : E:\seq_SU0411.txt
Output Set: N:\CRF4\04172006\J573885.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

Seq#:77; N Pos. 2199

see p. 4

10/573,885 4

SEQUENCE LISTING

change
to
↓
<110> SUNTORY LIMITED

<120> A gene encoding an enzyme for catalyzing biosynthesis of lignan,
and use thereof

<130> SU0411

<1507> PCT/JP2004/0146696
<1517> 2004-09-24

These are prior data

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/573,885

DATE: 04/17/2006
TIME: 12:27:25

Input Set : E:\seq_SU0411.txt
Output Set: N:\CRF4\04172006\J573885.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1812 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:77